

GAUAAUAUGGAUCUUUAAAGAUUCUUUUGUAAGCCCCAAGGGC (SEQ ID NO:22).

SUB  
E103  
108. (Amended) An RNA comprising the consensus sequence NNNGAUNCUUUNNGUAA GCCCNANGNGNN (SEQ ID NO:23) and having a first double stranded region, a first end loop region, a second double stranded region, and a second end loop region.

109. (Amended) An *in silico* representation of an RNA fragment that is conserved across at least two species comprising the sequence NNNGAUNCUUUNNGUAAAGCCCNANGNGNN (SEQ ID NO:23).

B<sup>4</sup>  
SUB  
F1  
110. (Amended) A purified and isolated RNA fragment that is conserved across at least two species comprising the sequence NNNGAUNCUUUNNGUAAAGCCCNANGNGNN (SEQ ID NO:23).

111. (Amended) A purified and isolated RNA fragment comprising the human sequence UAUGAUUCUUUUGUAAGCCCUAGGGGCU (SEQ ID NO:24).

112. (Amended) A purified and isolated RNA fragment comprising the mouse sequence AAAGAUUCUUUUGUAAGCCCCAAGGGCU (SEQ ID NO:25).

113. (Amended) A purified and isolated RNA fragment comprising the rat sequence AAAGAUUCUUUUGUAAGCCCCAAGGGCU (SEQ ID NO:25).

#### REMARKS

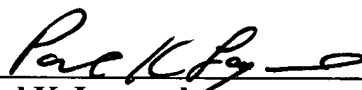
Claims 31-34, 56-59, 81-86 and 108-113 have been amended to incorporate sequence identifiers as requested by the Examiner in order to more fully comply with the Sequence Rules set forth in 37 CFR §§1.821-1.825. In addition, enclosed herewith is a copy of the Notice To Comply

**DOCKET NO.: IBIS-0171**

**PATENT**

With Requirements For Patent Applications Containing Nucleotide Sequence And/Or Amino Acid Sequence Disclosures and a response thereto. No new matter has been added. Applicants submit that the present response is complete.

Respectfully submitted,



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